

IN THE SPECIFICATION:

Please amend the paragraph beginning on page 27, line 31 as follows:

For the base sequence of K-ras oncogene, a database such as the GenBank was referenced, and probes (SEQ ID NO:56 to 62, 63 to 69, 70 to 76, 77 to 83) were solid-phased onto the microarray. In order to make the layout shown in FIG. 13, the composed probes were spotted using a micro dispersion system using a piezo device. At the spot 1 in FIG. 13, a 20mer sense strand probe corresponding to K-ras natural sequence (Wt) (SEQ ID NO:56) was arranged. At the spot 2, a 20mer sense strand probe corresponding to K-rasArg mutant (SEQ ID NO:57) was arranged. At the spot 3, a 20mer sense strand probe corresponding to K-rasCys mutant (SEQ ID NO:58) was arranged. At spots 4 to 7, 20mer sense strand probes corresponding to SEQ ID NOs 59-62, respectively were arranged. At the spot 8, a 20mer anti-sense strand probe corresponding to K-rasWt (SEQ ID NO:63) was arranged. At spots 9 to 14, 20mer anti-sense probes, corresponding to SEQ ID NOs 64-69 respectively, are arranged. At the spot 15, a 17mer sense strand probe corresponding to K-rasWt (SEQ ID NO:70) were arranged. At spots 16-21, 17mer sense probes, corresponding to SEQ ID NOs 71-76 respectively, were arranged. At the spot 22, a 17mer anti-sense strand probe corresponding to K-rasWt (SEQ ID NO:77) was arranged. At spots 23-28, 17mer anti-sense probes, corresponding to SEQ ID NOs 78-83, were arranged.